SEQUENCE LISTING

(1) GENERAL INFORMATION:

(i) APPLICANT: Cassels,, Frederick Lees, Andrew Schuman, Richard

- (ii) TITLE OF INVENTION: MONOCLONAL ANTIBODY WHICH AGGLUTINATES E. COLI HAVING THE CS4-CFA/1 FAMILY PROTEIN
- (iii) NUMBER OF SEQUENCES: 3
- (iv) CORRESPONDENCE ADDRESS:
 - (A) ADDRESSEE: Glenna Hendricks
 - (B) STREET: P.O. Box 2509
 - (C) CITY: Fairfax
 - (D) STATE: VA
 - (E) COUNTRY: USA
 - (F) ZIP: 22031
 - (v) COMPUTER READABLE FORM:
 - (A) MEDIUM TYPE: Floppy disk
 - (B) COMPUTER: IBM PC compatible
 - (C) OPERATING SYSTEM: PC-DOS/MS-DOS
 - (D) SOFTWARE: PatentIn Release #1.0, Version #1.25
- (vi) CURRENT APPLICATION DATA:
 - (A) APPLICATION NUMBER: US 08/905,046
 - (B) FILING DATE: 01-AUG-1997
 - (C) CLASSIFICATION:
- (viii) ATTORNEY/AGENT INFORMATION:
 - (A) NAME: Hendricks, Glenna
 - (B) REGISTRATION NUMBER: 32,535
 - (C) REFERENCE/DOCKET NUMBER: cassels/mon

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- (ix) TELECOMMUNICATION INFORMATION:
 - (A) TELEPHONE: (703) 591-4470
 - (B) TELEFAX: (703) 591-4428
- (2) INFORMATION FOR SEQ ID NO:1:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 36 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: unknown
 - (ii) MOLECULE TYPE: peptide
 - (iii) HYPOTHETICAL: NO

- (iv) ANTI-SENSE: NO
 - (v) FRAGMENT TYPE: internal
- (vi) ORIGINAL SOURCE:
 - (A) ORGANISM: E. COLI
 - (B) STRAIN: CS4-CFA/I
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

Val Glu Lys Asn Ile Thr Val Thr Ala Ser Val Asp Pro Thr Ile
Asp

1 5 10 15

Leu Leu Gln Ala Asp Gly Ser Ala Leu Pro Ser Ala Val Ala Leu Thr 20 25 30

Tyr Ser Pro Ala 35

- (2) INFORMATION FOR SEQ ID NO:2:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 8 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: unknown
 - (ii) MOLECULE TYPE: peptide
 - (iii) HYPOTHETICAL: NO
 - (iv) ANTI-SENSE: NO
 - (v) FRAGMENT TYPE: internal
 - (vi) ORIGINAL SOURCE:
 - (A) ORGANISM: E. COLI
 - (B) STRAIN: CS4-CFA/I
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

Ser Ala Val Ala Leu Thr Tyr Ser 1 5

- (2) INFORMATION FOR SEQ ID NO:3:
 - (i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 37 amino acids

- (B) TYPE: amino acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: unknown
- (ii) MOLECULE TYPE: peptide
- (iii) HYPOTHETICAL: NO
 - (iv) ANTI-SENSE: NO
 - (v) FRAGMENT TYPE: internal
 - (vi) ORIGINAL SOURCE:
 - (A) ORGANISM: E. COLI
 - (B) STRAIN: CS4-CFA/I
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:

Cys Val Glu Lys Asn Ile Thr Val Thr Ala Ser Val Asp Pro Thr Ile

1 5 10 15

Asp Leu Leu Gln Ala Asp Gly Ser Ala Leu Pro Ser Ala Val Ala Leu 20 25 30

Thr Tyr Ser Pro Ala 35

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